SEQUENCE LISTING

- (1) GENERAL INFORMATION
- (i) APPLICANT: Reich, Norbert O. Flynn, James
- (ii) TITLE OF THE INVENTION: MODULATORS OF DNA CYTOSINE-5 METHYLTRANSFERASE AND METHODS FOR USE THEREOF
- (iii) NUMBER OF SEQUENCES: 110
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Merchant, Gould, Smith, Edell, Welter & Schmidt (B) STREET: 11150 Santa Monica Boulevard, Suite 400

 - (C) CITY: Los Angeles
 - (D) STATE: CA
 - (E) COUNTRY: USA
 - (F) ZIF: 90025
- (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Diskette
 - (B) COMPUTER: IBM Compatible

 - (C) OPERATING SYSTEM: DOS
 (D) SOFTWARE: FastSEQ for Windows Version 2.0
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER:
 - (B) FILING DATE: 12-JUN-1998
 - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
- (A) APPLICATION NUMBER: 60/057,411
 - (B) FILING DATE: 29-AUG-1997
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Canady, Karen S
 - (B) REGISTRATION NUMBER: 39,927
 - (C) REFERENCE/DOCKET NUMBER: 30794.30W001
- (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: 310 445-1140 (B) TELEFAX: 310 445-9031

 - (C) TELEX:
 - (2) INFORMATION FOR SEO ID NO:1:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

GGGGGGRRK KGCGKGGKGK KGKKGG

- (2) INFORMATION FOR SEQ ID NO: 2:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 base pairs
 - (B) TYPE: nucleic acid (C) STRANDEDNESS: double

 - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:



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| | (D) TOPOLOGY: linear | ÷ |
| | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2: | |
| KGGRKI | KRDDD KRCGKRRDKK KKKKKG | 26 |
| | (2) INFORMATION FOR SEQ ID NO:3: | |
| | (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 10 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | · |
| | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3: | |
| GGGGCG | GGGGC | 10 |
| | (2) INFORMATION FOR SEQ ID NO:4: | • |
| | (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 10 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4: | |
| CCAACG | TTGG | 10 |
| | (2) INFORMATION FOR SEQ ID NO:5: | |
| | (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 10 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5: | |
| CCAAGA! | TTGG | 10 |
| | (2) INFORMATION FOR SEQ ID NO:6: | |

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 10 base pairs (B) TYPE: nucleic acid

(C) STRANDEDNESS: double

(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

CGATCGATCG

(2) INFORMATION FOR SEQ ID NO:7:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 10 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: double

 - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

| CCAACGTTGG | 10 |
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| (2) INFORMATION FOR SEQ ID NO:8: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ 10 NO:8: | |
| CCGTACGTAC GG | 12 |
| (2) INFORMATION FOR SEQ ID NO:9: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (x1) SEQUENCE DESCRIPTION: SEQ ID NO:9: | |
| GGGAATTCAA GGGGCGGGC AAGGATCCAG | 30 |
| (2) INFORMATION FOR SEQ ID NO:10: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | · · · · · . |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: | |
| CTGGATCCTT GCCCCGCCCC TTGAATTCCC | 30 |
| (2) INFORMATION FOR SEQ ID NO:11: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: | |
| GGGAATTCAA ATGACGTCAA AAGGATCCAG | 30 |
| (2) INFORMATION FOR SEQ ID NO:12: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:12: | |
| CTGGATCCTT TTGACGTCAT TTGAATTCCC | 30 |
| (2) INFORMATION FOR SEC ID NO. 13. | |



| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 50 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
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| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:13: | |
| CCTACCCACC CTGGATCCTT GCCCCGCCCC TTGAATTCCC AACCCTCCAC | 50 |
| (2) INFORMATION FOR SEQ ID NO:14: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 22 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:14: | |
| ATCCTTGCCC CGCCCCTTGA AT | 22 |
| (2) INFORMATION FOR SEQ ID NO:15: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 14 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (x1) SEQUENCE DESCRIPTION: -SEQ-ID-NO: 15: | |
| TTGCCCCGCC CCTT | 14 |
| (2) INFORMATION FOR SEQ ID NO:16: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 66 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:16: | |
| GGGAATTCAT GGATCCTAAA NNNNNNNNN NNCGNNNNNN NNNNNNTTTC AAGCTTGTGA | 60 66 |
| (2) INFORMATION FOR SEQ ID NO:17: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 66 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:17: | |
| CCCTTAAGTA CCTAGGATTT NNNNNNNNN NNGCNNNNNN NNNNNNAAAG TTCGAACACT TAAGGG | 60 66 |
| (2) INFORMATION FOR SEQ ID NO:18: | |
| (i) CROURNOR OURDACHERYCON | |

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| (A) LENGTH: 26 base pairs | |
| (B) TYPE: nucleic acid | |
| (C) STRANDEDNESS: double | |
| (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:18: | |
| GTGGGATGGG AACGAGTTGA GGAGGG | 2 |
| (2) INFORMATION FOR SEQ ID NO:19: | |
| (i) SEQUENCE CHARACTERISTICS: | |
| (A) LENGTH: 26 base pairs | |
| (B) TYPE: nucleic acid | |
| (C) STRANDEDNESS: double | |
| (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:19: | |
| AGTGGTATGT ATCGATTATA GTTGGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:20: | |
| (i) SEQUENCE CHARACTERISTICS: | |
| (A) LENGTH: 26 base pairs | |
| (B) TYPE: nucleic acid | |
| (C) STRANDEDNESS: double | |
| (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:20: | |
| GGAGGAAGTT TACGTATGGT ATGGGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:21: | 26 |
| | |
| (i) SEQUENCE CHARACTERISTICS: | |
| (A) LENGTH: 26 base pairs | |

- (B) TYPE: nucleic acid(C) STRANDEDNESS: double
- (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:

TGGGAGGGA TTCGAGGTGA GAGTTG

- (2) INFORMATION FOR SEQ ID NO:22:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: double

 - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:

ATAAAGTATT AGCGTAAGAG ATGAAG

26

- (2) INFORMATION FOR SEQ ID NO:23:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 base pairs

 - (B) TYPE: nucleic acid
 (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear



| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:23: | |
|--|-----|
| TGGAGGAGTT TACGGTGTAA TTGTTT | 26 |
| (2) INFORMATION FOR SEQ ID NO:24: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ 1D NO:24: | |
| GGAGTAGGTA GACGTTAAGT ATGATG | 2 б |
| (2) INFORMATION FOR SEQ ID NO:25: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:25: | |
| GTGGGAAGGG GACGAATTTG AAGGTG | 26 |
| (2) INFORMATION FOR SEQ ID NO:26: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | - |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:26: | |
| TGGTAATGTA TTCGTAAATG TAAGGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:27: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:27: | |
| TAATAGGGGA GACGTAAATG TAAGGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:28: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:28: | |

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| GAGTGTAGAA GTCGTAATAG ATTTAG | 26 |
|--|--------------|
| (2) INFORMATION FOR SEQ ID NO:29: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:29: | |
| TGAGTAGGAA AGCGAAGAGG TGTTGG | 26 |
| (2) INFORMATION FOR SEQ ID NO: 30: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:30: | |
| TAGGTATTGG GGCGGAAGGT GGGTGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:31: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | - |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:31: | |
| GGGGGTATAA TACGGTGTTG GTAGGG | 26 |
| (2) INFORMATION FOR SEQ ID NO: 32: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:32: | |
| GGGTTGGGGT TTCGTGTGGG GGGTGT | 26 |
| (2) INFORMATION FOR SEQ ID NO:33: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:33: | |
| TGTGGGTATG GGCGGTGATA GTGAAG | . 26 |
| 12) THEORY TON TON TO 10 10 12. | |



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- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 base pairs(B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:34:

GGATGATGGG GTCGAGAGTG GTGGTG

26

- (2) INFORMATION FOR SEQ ID NO:35:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:35:

TAGTGGGTGG AGCGAGTGGT GGTTGG

26

- (2) INFORMATION FOR SEQ ID NO: 36:
- - (i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 26 base pairs

 - (B) TYPE: nucleic acid (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:36:

AGGGTGGGTG GGCGGAGTTG TTGTTG

26

- (2) INFORMATION FOR SEQ ID NO: 37:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 37:

GTGAGGAGGG AGCGGGAATG GGGGTG

26

- (2) INFORMATION FOR SEQ ID NO:38:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:38:

GGGGGTGGG AGCGGAGGGG GGTGAG

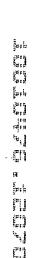
26

- (2) INFORMATION FOR SEQ ID NO: 39:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid

 - (C) STRANDEDNESS: double

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| (D) TOPOLOGY: linear | |
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| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:39: | |
| TGTTGGAGGG GGCGAAGGTG GTTTTG | 2 |
| (2) INFORMATION FOR SEQ ID NO:40: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | · |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 40: | |
| GGGGGGGGG GGCGAGGGGT AGATGG | 2: |
| (2) INFORMATION FOR SEQ ID NO:41: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:41: | 1 |
| GGGGGAGGG TTCGGTGATA GGTAGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:42: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:42: | |
| GGGGGGGGG TACGTGGGAT GGTATG | 26 |
| (2) INFORMATION FOR SEQ ID NO: 43: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | , |
| | |

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:43:

GTGTAGGGG TGTAAA

- (2) INFORMATION FOR SEQ ID NO:44:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: double
 (D) TOPOLOGY: linear
- (x1) SEQUENCE DESCRIPTION: SEQ ID NO:44:



| GGGGGGGGT AGCGCTTAGA TGGTGG | 26 |
|--|----|
| (2) INFORMATION FOR SEQ ID NO:45: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:45: | |
| GGGGTGAGGG GGCGGGGGTT AGTGGG | 26 |
| (2) INFORMATION FOR SEQ ID NO: 46: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:46: | |
| GAGGGGGGT TGCGTAGGGG GGTGGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:47: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:47: | |
| TGTGGAGGTG GGCGGGAAAG GTGATG | 26 |
| (2) INFORMATION FOR SEQ ID NO:48: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:48: | |
| CGGGGGATGG GACGGATGGG GGGGGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:49: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:49: | |
| GGGGGTGGGG TGCGAGAGAG TTGGGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:50: | |

| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
|--|----|
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:50: | |
| GAGGGGTGGA GGCGGAGGTG GGTTGG | 2 |
| (2) INFORMATION FOR SEQ ID NO:51: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:51: | • |
| GGGGGGGGG GGCGGATAAG GGTGTG | 26 |
| (2) INFORMATION FOR SEQ ID NO:52: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:52: | |
| TGGGGGGGG GGCGGGGGA GTTTGA . | 26 |
| (2) INFORMATION FOR SEQ ID NO:53: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:53: | |
| GGGGGGGAGG GGCGGATAGT TGTGTG | 26 |
| (2) INFORMATION FOR SEQ ID NO:54: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | · |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:54: | |
| GGGTGGGGTG GGCGGTGGGG TGTGGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:55: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid | |

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| <pre>(C) STRANDEDNESS: double (D) TOPOLOGY: linear</pre> | |
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| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:55: | |
| GAGGGGGGG AGCGGAGGGG GTTGGG | 2 |
| (2) INFORMATION FOR SEQ ID NO:56: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:56: | |
| GGGGGGAAG GGCGTGGGGT TGGGTG | 2€ |
| (2) INFORMATION FOR SEQ ID NO:57: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:57: | |
| GCGAGGGGG GCGATGGGGT GGTGG | 25 |
| (2) INFORMATION FOR SEQ ID NO:58: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:58: | |
| GGGTGGGGGT GGCGTTGTGG GTGGGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:59: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:59: | |

(2) INFORMATION FOR SEQ ID NO:60:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 26 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: double
(D) TOPOLOGY: linear

GGGAGGGGT GGCGGTGGGT ATGTGG

| ANTA DEGOEAGE DESCRIPTION: SEQ 10 NO: 60: | |
|--|-----|
| GGGGAGGGTG GGCGGGTATG GAGTGG | 2 |
| (2) INFORMATION FOR SEQ ID NO:61: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:61: | |
| GGGGGGGAG TGCGTTGATG GGTGTG | 26 |
| (2) INFORMATION FOR SEQ ID NO: 62: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:62: | |
| GGGGGGTGG ATCGTGGGGG GAGGGG | 26 |
| . (2) INFORMATION FOR SEQ ID NO:63: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:63: | |
| GGGGTAGGGT GGCGGGGGG GTATGG | 26 |
| (2) INFORMATION FOR SEQ ID NO: 64: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:64: | |
| GGGATGGGGG TGCGGGGTAT GGGGGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:65: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:65: | |
| GGGAGGGGT AGCGGGAGTG TGTGTG | 0.0 |

(2) INFORMATION FOR SEQ ID NO:66:

| PCT/US98/12351 |
|----------------|

| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
|--|----|
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:66: | |
| GGGGCTAAGG GGCGTAAGAA TGGGGG | 2 |
| (2) INFORMATION FOR SEQ ID NO:67: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:67: | |
| GGGGGGGTGG TTCGGTAATG GGGGGT | 26 |
| (2) INFORMATION FOR SEQ ID NO:68: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:68: | |
| GGTGGGAGAG GGCGTGGTGT AGGTAG | 26 |
| (2) INFORMATION FOR SEQ ID NO:69: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:69: | |
| GGGGGGGTG TACGAGGTTT GTGTGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:70: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | : |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:70: | |
| TGGTGGAGGG GGCGAAGAAG TGTGTG | 26 |

(2) INFORMATION FOR SEQ ID NO:71:

(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs

| (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
|--|----|
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:71: | |
| GGGGGTGGGA TGCGGAATAA GGATGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:72: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:72: | |
| TGAGGGGGAG GGCGAATAGA TGGTGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:73: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:73: | |
| GGGGGGAGTA AGCGGGGGTG TGGTGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:74: | - |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:74: | |
| TGAAGGGGGG TGCGGGGTGT GGGGG | 25 |
| (2) INFORMATION FOR SEQ ID NO:75: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (wi) PEOUPLOS DECONTRATON AND AN AND AND AND AND AND AND AND AND | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:75: | |
| GTGGTGATGG GGCGGGGTGG TAGTGG | 26 |

- (2) INFORMATION FOR SEQ ID NO:76:
- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 26 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: double
 (D) TOPOLOGY: linear

| (XI) SEQUENCE DESCRIPTION: SEQ ID NO: /6: | |
|--|-----|
| TGGAGGGGTA GGCGTGGGGT GATGGG | 2 |
| (2) INFORMATION FOR SEQ ID NO:77: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:77: | |
| GGTAGGGAGT GGCGGGTGGT GATGGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:78: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:78: | |
| GGGTGTAGAG GGCGGGAGTA GAGGGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:79: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | - |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:79: | |
| GGGTGGGTTT GGCGTAATTG TGTGGG | 2.6 |
| (2) INFORMATION FOR SEQ ID NO:80: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:80: | |
| GGGTGTGTTG GGCGTGGGGT ATGTAG | 26 |
| (2) INFORMATION FOR SEQ ID NO:81: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:81: | |
| TGGGGAGAAT GGCGGGGGT GGTGGG | 26 |





(2) INFORMATION FOR SEQ ID NO:82:

| (i) | SEQUENCE | CHARACTERISTICS: |
|-----|----------|--------------------|
| / | 22502465 | CHUINCI PUTDITICS: |

- (A) LENGTH: 26 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:82:

TATGGTGGGA GGCGGGGGGG GGTTGG

26

(2) INFORMATION FOR SEQ ID NO:83:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:83:

TGGGGAAAGA GGCGTGAGTG GGGGGG

26

(2) INFORMATION FOR SEQ ID NO:84:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:84:

TGTAGGGGAG GACGGGGGAT GGGGTG

26

(2) INFORMATION FOR SEQ ID NO:85:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:85:

GGGTGGGTAA TGCGTAGGGT GGGGGG

26

(2) INFORMATION FOR SEQ ID NO:86:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:86:

GTGTGGGTAA GGCGGTATGG GGGTGG

26

(2) INFORMATION FOR SEQ ID NO:87:

(i) SEQUENCE CHARACTERISTICS:

| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:87: | |
|--|-----|
| TGGAGGGTGT TGCGGTGAGG TGGTGG | 2 € |
| (2) INFORMATION FOR SEQ ID NO:88: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:88: | |
| GGTGGTGGTG ATCGGGGTTG TGATGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:89: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:89: | |
| GGGGGTAAAG TGCGGGTGGT TGATGG | 26 |
| (2) INFORMATION FOR SEQ ID NO: 90: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:90: | |
| GTGGAGGTGT TGCGTAGTGT GGGAGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:91: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:91: | |
| STGGGGAATG GTCGGTTATG GTGGGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:92: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double | |

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:92:

| GGGATGTGGT AGCGGGGGTG TGTTAG | 2 |
|---|-----------|
| (2) INFORMATION FOR SEQ ID NO:93: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:93: | |
| GGGGTAGGAG TTCGTAGGGG TGTGTT | 26 |
| (2) INFORMATION FOR SEQ ID NO:94: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:94: | |
| GAGGTGGTGG ATCGGGATGA TGGATT | 26 |
| (2) INFORMATION FOR SEQ ID NO:95: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | *** ***** |
| · | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:95: | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:95: TGGGGGGAAA TACGGGGAGG GTGGTA | 26 |
| | 26 |
| TGGGGGGAAA TACGGGGAGG GTGGTA | 26 |
| TGGGGGGAAA TACGGGGAGG GTGGTA (2) INFORMATION FOR SEQ ID NO:96: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double | 26 |
| TGGGGGGAAA TACGGGGAGG GTGGTA (2) INFORMATION FOR SEQ ID NO:96: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | 26 |
| TGGGGGGAAA TACGGGGAGG GTGGTA (2) INFORMATION FOR SEQ ID NO:96: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear (xi) SEQUENCE DESCRIPTION: SEQ ID NO:96: | |
| TGGGGGGAAA TACGGGGAGG GTGGTA (2) INFORMATION FOR SEQ ID NO:96: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear (xi) SEQUENCE DESCRIPTION: SEQ ID NO:96: GGAGTAGGGT TACGTGGTGG TAATGG | |
| TGGGGGGAAA TACGGGGAGG GTGGTA (2) INFORMATION FOR SEQ ID NO:96: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear (xi) SEQUENCE DESCRIPTION: SEQ ID NO:96: GGAGTAGGGT TACGTGGTGG TAATGG (2) INFORMATION FOR SEQ ID NO:97: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double | |

(2) INFORMATION FOR SEQ ID NO:98:

| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
|--|------|
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:98: | |
| TGGATGAGAG TGCGTGTATG ATAAGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:99: | • |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:99: | |
| AGGCTTAGTG AACGGGGGG AGGTGG | 26 |
| (2) INFORMATION FOR SEQ ID NO:100: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:100: | |
| GAGAAGGGTA AACGTGGGGG AGGGGA | - 26 |
| (2) INFORMATION FOR SEQ ID NO:101: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:101: | |
| TGGGGGGGG GGCGGGGGA GTTTGA | 26 |
| (2) INFORMATION FOR SEQ ID NO:102: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 56 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (5) Islands. Ithear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:102: | |
| GAACAATGGG GGCGCTGGGG GGGGGGCGGGGCTTTA GCTATGTCAG AATTCA | 56 |
| (2) INFORMATION FOR SEQ ID NO:103: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 bas pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double | - |
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| (xi) | SEQUENCE DESCRIPTION: SEQ ID NO:103: |
|------------|--------------------------------------|
| GGGATGGGGG | TGCGGGGTAT GGGGGG |
| (2 | 2) INFORMATION FOR SEC ID NO.104. |

(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 57 base pairs (B) TYPE: nucleic acid

(D) TOPOLOGY: linear

- (C) STRANDEDNESS: double (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:104:

GGGGAACAGC GAGCACCGAA GGGGGTGCGG GGTATGGGAG GGTCCCCGGG CTTGAGC 57

(2) INFORMATION FOR SEQ ID NO: 105:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:105:

GGTGGTGGTG ATCGGGGTTG TGATGG

(2) INFORMATION FOR SEQ ID NO: 106:

- (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 56 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 106:

TGTCCTTCTT GTGGTGGTGG TAGAGGTCGT GGTTGTGATG GTGGCTCGGT GTGTGT 56

- (2) INFORMATION FOR SEQ ID NO:107:
- (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs

 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear
- (x1) SEQUENCE DESCRIPTION: SEQ ID NO:107:

GAGGGGGG AGCGGAGGG GTTGGG

26

- (2) INFORMATION FOR SEQ ID NO:108:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 56 base pairs (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:108:

| -81- | |
|--|----|
| GAGGGGTGTA GCCGCGAGGG GGCGGAGCGG AGGGGGAGGG CCCTGGTCCC GCCGCC | 56 |
| (2) INFORMATION FOR SEQ ID NO:109: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:109: | |
| CCCCACCCAC AACGCCACCC CCACCC | 26 |
| (2) INFORMATION FOR SEQ ID NO:110: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 56 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear | |

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:110: TCTTTAAATG GTGCGGTCCA CCCCCACCGC CACCCCCACC CCCCACTGGA GCAAGG 56

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